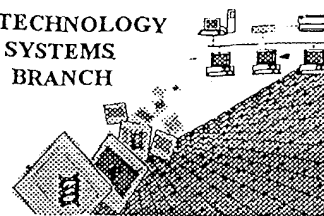


RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/974,619C
Source: 1600
Date Processed by STIC: 10/17/2003

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

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<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

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Revised 10/08/2003



1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/974,619C

DATE: 10/17/2003

TIME: 14:49:23

Input Set : A:\sj29sqlist.txt

Output Set: N:\CRF4\10172003\I974619C.raw

4 <110> APPLICANT: Schuetz, Erin
 5 Zhang, Joing
 6 Assem, Mahfoud
 8 <120> TITLE OF INVENTION: Genotyping Assay to Predict CYP3A5
 9 Phenotype
 11 <130> FILE REFERENCE: 44158/244344
 13 <140> CURRENT APPLICATION NUMBER: 09/974,619C
 14 <141> CURRENT FILING DATE: 2001-10-10
 16 <150> PRIOR APPLICATION NUMBER: 60/279,915
 17 <151> PRIOR FILING DATE: 2001-03-29
 19 <160> NUMBER OF SEQ ID NOS: 70
 21 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

563 <210> SEQ ID NO: 49
 564 <211> LENGTH: 31 14 shown below
 565 <212> TYPE: DNA
 566 <213> ORGANISM: Homo sapien
 568 <400> SEQUENCE: 49
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 752 <210> SEQ ID NO: 70
 753 <211> LENGTH: 13
 754 <212> TYPE: DNA
 755 <213> ORGANISM: Homo sapien
 757 <400> SEQUENCE: 70
 758 tattgtagat ccc
 E--> 762 1
delete

Does Not Comply
Corrected Discrete Needed

14

13

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/974,619C

DATE: 10/17/2003

TIME: 14:49:24

Input Set : A:\sj29sqlist.txt

Output Set: N:\CRF4\10172003\I974619C.raw

L:569 M:252 E: No. of Seq. differs, <211> LENGTH:Input:31 Found:14 SEQ:49

L:762 M:254 E: No. of Bases conflict, this line has no nucleotides.